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Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

- 1. (currently amended) An isolated DNA construct comprising at least one mutated binding site for a growth factor independence–1 (Gfi-1) transcription repressor, said mutated binding site comprising a <u>point</u> mutation which hinders or prevents binding of said Gfi-1 repressor to said site.
 - 2. (original) The DNA construct of claim 1, which is a promoter.
- 3. (original) The DNA construct of claim 2, wherein said promoter is a mammalian cellular promoter.
- 4. (original) The DNA construct of claim 2, wherein said promoter is a viral promoter.
- 5. (original) The DNA construct of claim 4, wherein said promoter is a human cytomegalovirus promoter.
- 6. (currently amended) The DNA construct of claim 5, which is a cytomegalovirus MIE (<u>Major Immediate Early</u>) promoter.
- 7. (previously presented) The DNA construct of claim 1, wherein said Gfi-1 binding site prior to said mutation is greater than 65% homologous with a sequence consisting of TAAATCACNGCA (Sequence I.D. No. 2), wherein N is A or T.
- 8. (previously presented) The DNA construct of claim 1, wherein said Gfi-1 binding site prior to said mutation is greater than 79% homologous with a sequence consisting of

TAAATCACNGCA (Sequence I.D. No. 2), wherein N is A or T.

- 9. (original) The DNA construct of claim 1, wherein said Gfi-1 binding site prior to said mutation comprises the sequence $N_1AAATCACN_2GCA$ (Sequence I.D. No. 1), wherein N_1 and N_2 are any nucleotide, and said mutation is in a portion of said binding site comprising the sequence AATC.
- 10. (original) The DNA construct of claim 1, wherein said binding site resides within an expression regulatory segment and said regulatory segment is operatively linked to a coding segment.
- 11. (original) The DNA construct of claim 10, wherein the coding segment encodes a gene product selected from the group consisting of cytokines, interleukins, interferons, growth factors and proto-oncogenes.
- 12. (withdrawn) An expression regulatory segment comprising at least one copy of a sequence N₁A-R-CN₂AGCA (Sequence I.D. No. 3), wherein N₁ and N₂ are any nucleotide, and R is a tetranucleotide selected from the group consisting of:

N₃ATC, AN₄TC, AAN₅C, AATN₆

N₃N₄TC, N₃AN₅C, N₃ATN₆, AN₄N₅C, AN₄TN₆, AAN₅N₆

N₃N₄N₅C, N₃N₄TN₆, N₃AN₅N₆, AN₄N₅N₆, and N₃N₄N₅N₆,

wherein N₃ is G, C or T, or is absent, or is an oligonucleotide of two or more nucleotides;

N₄ is G, C or T, or is absent, or is an oligonucleotide of two or more nucleotides;

N₅ is A, G or C, or is absent, or is an oligonucleotide of two or more nucleotides; and

N₆ is A, G or C, or is absent, or is an oligonucleotide of two or more nucleotides.

13. (withdrawn) The expression regulatory segment of claim 12, wherein R is selected from the group consisting of GATC, ACTC and AATA.

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14. (withdrawn) The expression regulatory segment of claim 12, which is a promoter.

- 15. (withdrawn) The expression regulatory segment of claim 14, wherein said promoter is a mammalian cellular promoter.
- 16. (withdrawn) The expression regulatory segment of claim 14, wherein said promoter is a viral promoter.
- 17. (withdrawn) The expression regulatory segment of claim 16, wherein said promoter is a human cytomegalovirus promoter.
- 18. (withdrawn) The expression regulatory segment of claim 17, which is a human cytomegalovirus MIE promoter.
- 19. (withdrawn) An expression vector comprising the expression regulatory segment of claim 12 and an operatively positioned insertion site for insertion of a coding segment.
- 20. (withdrawn) The expression vector of claim 19, in which is inserted a coding segment selected from the group consisting of cytokines, interleukins, interferons, growth factors and proto-oncogenes.
- 21. (original) An isolated DNA molecule comprising a sequence selected from the group consisting of Sequence I.D. No. 13 and Sequence I.D. No. 14.
 - 22. (original) An expression vector comprising the DNA molecule of claim 21.
- 23. (original) A method for improving expression of genes regulated by expression regulatory sequences which contain binding sites for a Gfi-1 transcription repressor,

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which comprises altering the sequence of said binding sites so as to hinder or prevent binding of said Gfi-1 transcription repressor to said binding sites, thereby improving said gene expression.

24. (currently amended) The method of claim 23, wherein said binding sites are altered at a tetranucleotide sequence contained therein, which wherein the tetranucleotide sequence is AATC.